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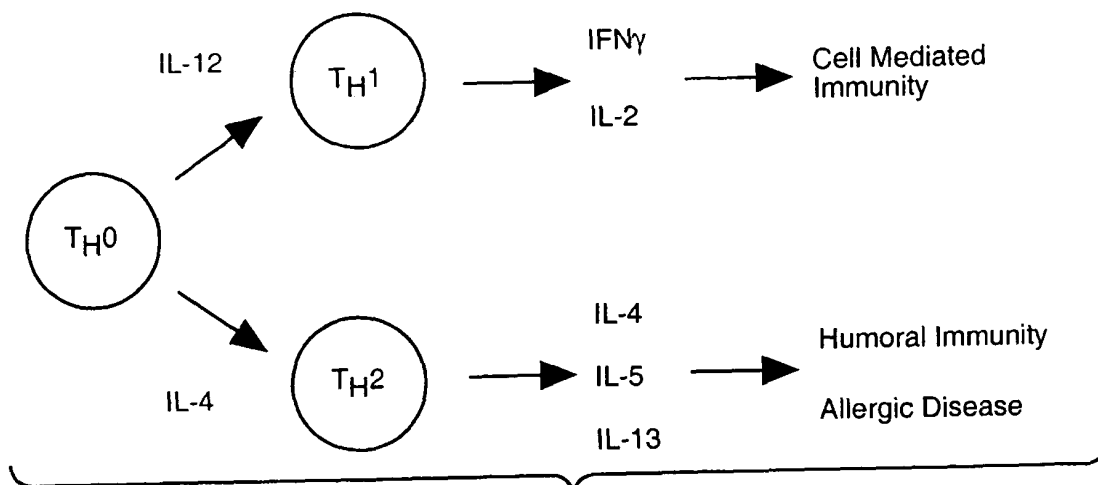


FIG. 1

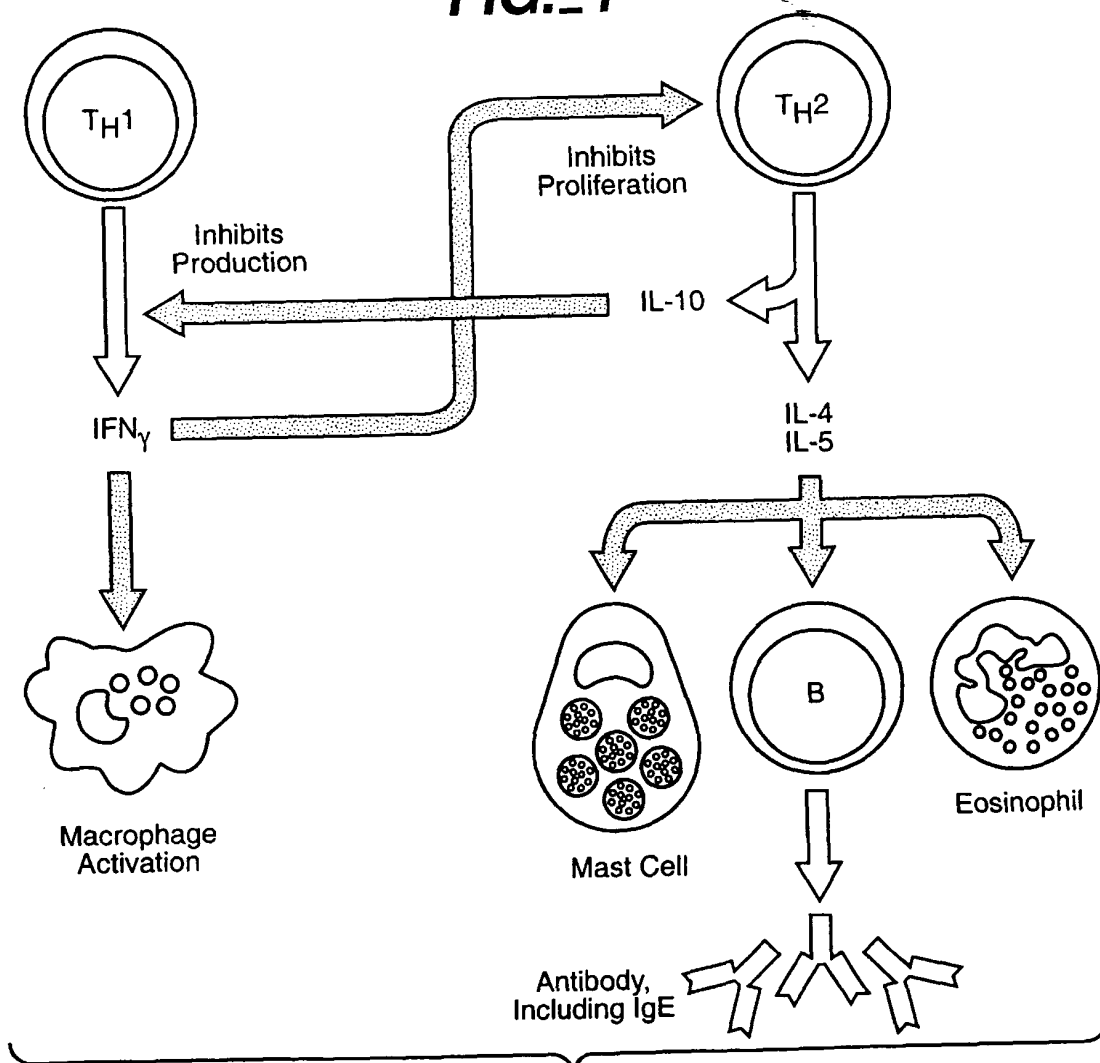


FIG. 2

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MRGGRGAPFWLWPLPKLALLPLLWVLFQRTRPQGSAGPLQCYGVGPLGDLNCSWEPLGD
LGAPSELHLQSQKYRSNKTQTVAVAAGRSWVAIPREQLTMSDKLLVWGTKAGQPLWPPV
FVNLETQMKPNAPRLGPDVDFSEDDPLEATVHWAPP TWP SHKVLICQFHYRRCQEAAWT
LLEPELKTIP LTPVEIQDLELATGYKVYGRCRMEKEEDLWGEWSPILSFQTPPSAPKDV
WVSGNLCGTPGGEELLLLWKAPGPCVQVSYKVWFWVGGRELSPEGITCCCSLIPSGAEW
ARVSAVNATSWEPLTNLSLVCLDSASAPRSVAVSSIAGSTELLVTWQPGPGEPLHVVD
WARDGDPLEKLNWVRLPPGNLSALLPGNFTVGVPRITVTAVSASGLASASSVWGFREE
LAPLVGPTLWRLQDAPPGTPAIAWGEVPRHQLRGHLTHYTLCAQSGTSPSVC MNVSGNT
QSVTLPDLPWGPCELWVTASTIAGQGPPGPILRLHLPDNTLRWKVLP GILFLWGLFLLG
CGLSLATSGRCYHLRHKVLPRWVWEKVPDPANSSSGQPHMEQVPEAQPLGDLPILEVEE
MEPPPVMESSQPAQATAPLDSGYEKHFLPTPEELGLLGPPRPQVLA

FIG._3

MNRLRVARLTPLELLLSLMSLLLGTRPHGSPGPLQCYSVGPLGILNCSWEPLGDLETPPV
LYHQSQKYHPNRVWEVKVPSKQSWVTIPREQFTMADKLLIWGTQKGRPLWSSVSVNLETQ
MKPDTPQIFSQVDISEEATLEATVQWAPPVWPPQKALTCQFRYKECQAEAWTRLEPQLKT
DGLTPVEMQNLEPGTCYQVSGRCQVENGYPWGEWSSPLSFQTPFLDPEDVWVSGTVCETS
GKRAALLVWKDPRPCVQVTTYTVWFGAGDITTTQEEVPCKSPVPAWMEWAVVSPGNSTSW
VPPTNLSLVCLAPESAPCDVGVSADGSPGIKVTWKQGTTRKPLEYVVDWAQDGDSDLKLN
WTRLPPGNLSTLLPGEFKGVPYRITVTAVYSGGLAAAPSVWGFREELVPLAGPAVWRLP
DDPPGTPVVAWGEVPRHQLRGQATHYTFICIQSRGLSTVC RNVSSQTQTATLPNLHSGSFK
LWVTVSTVAGQGPPGPDLSLHLPDNRI RWKALPWFLSLWGLLLMGCGLSLASTRCLQARC
LHWRHKLLPQWIWERVDPANSNSGQPYIKEVSLPQPPKDGPILEVEEVELQPVVESPKA
SAPIYSGYEKHFLPTPEELGLLV

FIG._4

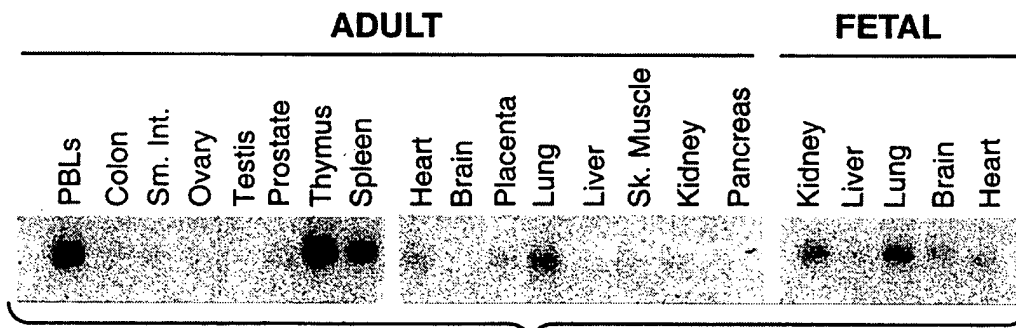


FIG._6

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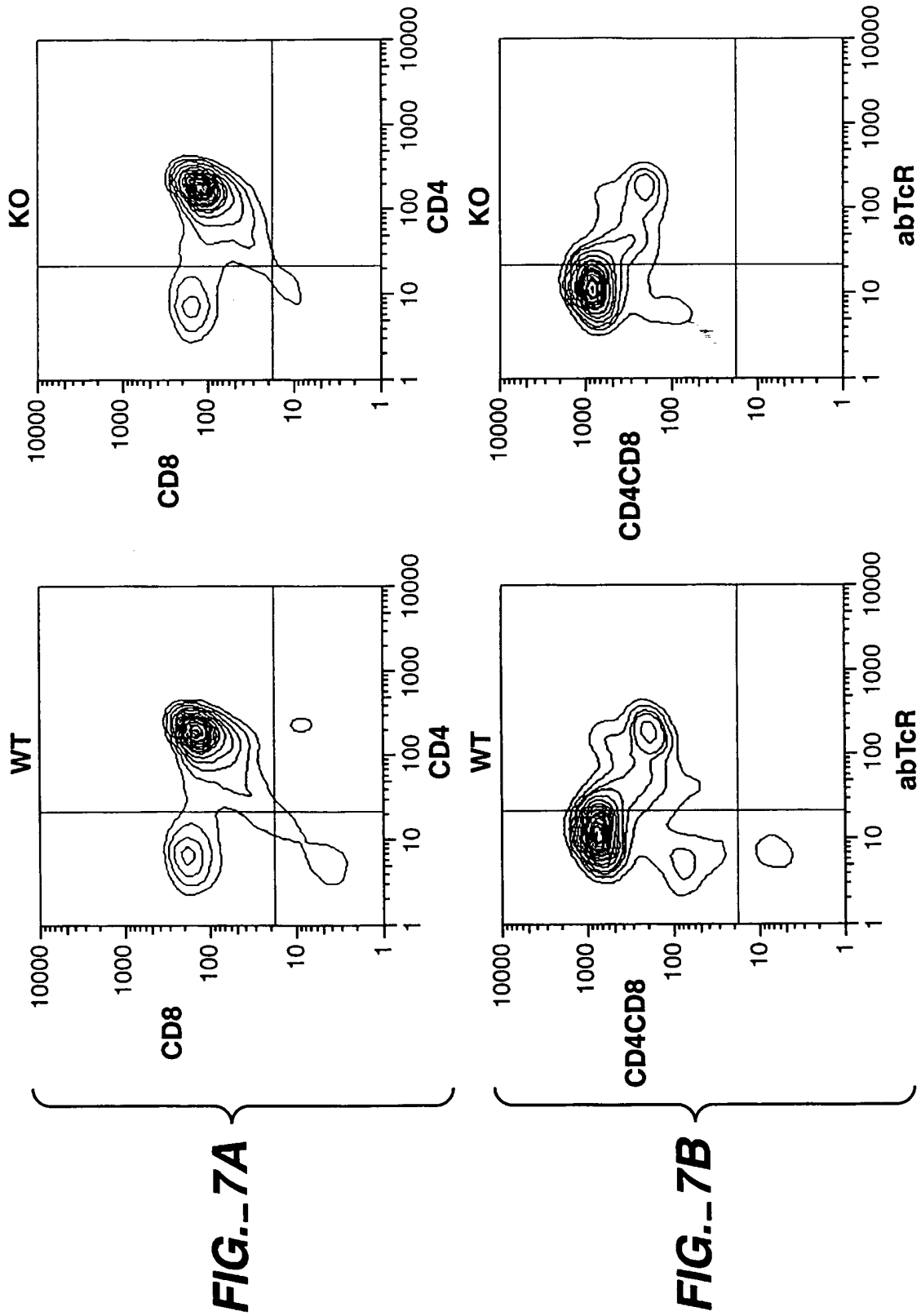
h-TCCR	1	MRGGRGAPFWLWPF	PKLALLPLLLVVF	FRTRPQGS	AGPLQCYG	VGPLGLDL
m-TCCR	1	-----MNRLRVAR	LTPLLELLLS	LLGTRPHG	SPGLQCY	SVGPLGIL
h-TCCR	51	NCSWEPLGDL	GAPSEELHLSQ	KYRSNKTQT	VAAAGR	SWVAIPREQLTMS
m-TCCR	46	NCSWEPLGDL	ETPPVLYHQ	SQKYHPNRV	WEVKVPSK	QSWVTIPREQFTMA
h-TCCR	101	DKLLVWGTKA	QQPLWPPV	VFNLE	TQMKPN	ARLGPDDVDFSEDDP
m-TCCR	96	DKLLVWGTKA	QCRPLWSS	SVSNLE	TQMKPD	TQIFSQVDISEATLEATVQ
h-TCCR	151	WAPPTWPSHR	VLCQFHYRR	CQEAATL	LEPELKT	IPLTPEIQQDLELAT
m-TCCR	146	WAPPVWPPQ	KALTCQFRYK	CCQAEATRL	LEPQLKT	DGLTPVEMQNLEPGT
h-TCCR	201	GKYYGRCRM	EEEDLWGEWS	PIESFQ	TPPSA	PKDVWVSGNLCTGTPGEE
m-TCCR	195	CYQVSGRCQ	VENGYP-WGEWS	PSLSFQ	TPFLD	PEBVDVWVGTVCE
h-TCCR	251	PLLLWKA	PGPCVQVS	YKVFVVG	GRELS	PEGITCCCSLI
m-TCCR	245	ALLVWKD	PRPCVQVT	TVWFG	AGDITTTQ	EEVPCCKSPVPAWMEAVVSP
h-TCCR	301	VNATSWEP	LTNLSLVCL	DSASAP	RSVA	VSSIAGSTELLVTWQPGGEPLE
m-TCCR	295	GMSVSWV	PPTNLSLVCL	APESAP	CDVGVSS	ADGSPGIKVTWKQGT
h-TCCR	351	HVVDWAR	DGDPLEKLN	VWRLP	PPGNLS	ALLPGNFTVGVPRITVTAVSASG
m-TCCR	345	YVVDWAQ	DGDSLDKLN	WTRLR	PPGNLS	TLLPGEFKGGVPRITVTAVYSGG
h-TCCR	401	LAASVSWG	FREELAPLV	GPTLLWRL	QDAPP	GTPAIAWGEVPRRHQLRGHLT
m-TCCR	395	LAAAPSVW	GFEELVPLA	GPAVWRL	PDDBP	PGTPVVAWGEVPRRHQLRGQAT
h-TCCR	451	HYTLCAQ	SGTSPSCMNV	SGNTQSV	TLPDLP	WGPCELWVTASTIAGQGPP
m-TCCR	445	HYTFCIQ	SRGLSTVCR	NVSSQTQ	TATLP	NLHSGSFKLWVTSTVAGQGPP
h-TCCR	501	GPILRLHLP	DNTRLRWK	VLPGIL	FLWGLF	LLGCGLSLATS-----GRCYHLR
m-TCCR	495	GPDLSLHLP	DNRIIRWK	ALPWF	LSLWGLLL	LMGCGLSLATS
h-TCCR	547	HKVLP	PRVWEK	VPDPANS	SSGQPH	MEQVPEAQPLGDLPILEVEEMEP
m-TCCR	545	HKLLPQW	IWERVP	DPANS	NSGQPY	IKEVSLPQPPKDGPILEVEEVELQPV
h-TCCR	597	MESSQPAQA	TAPLDDSG	YKHF	FLPTPE	ELGLLGGPPRPQVLA
m-TCCR	595	VES-----PKAS	APIYSG	YKHF	FLPTPE	ELGLLV

Box 1

FIG.-5

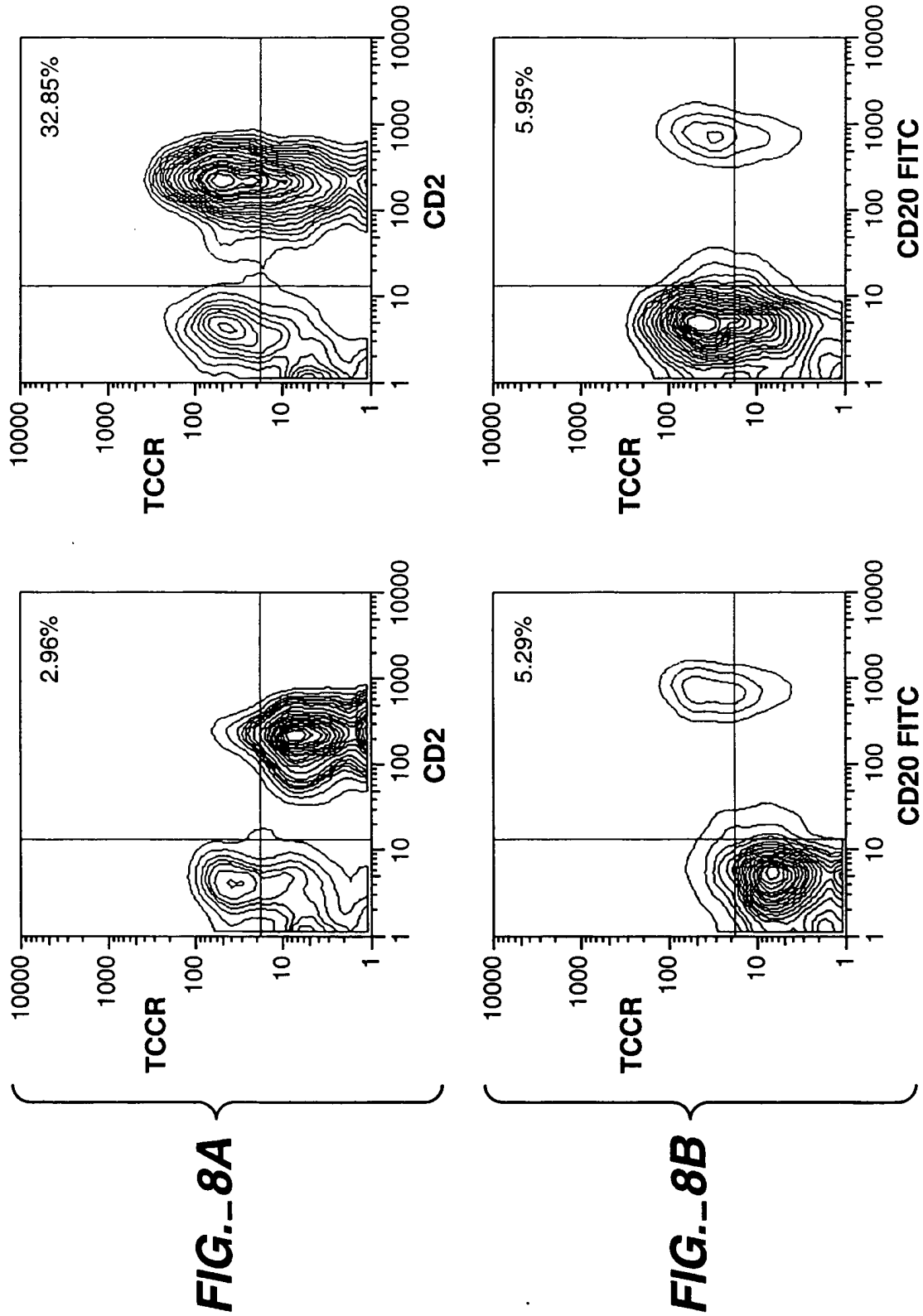
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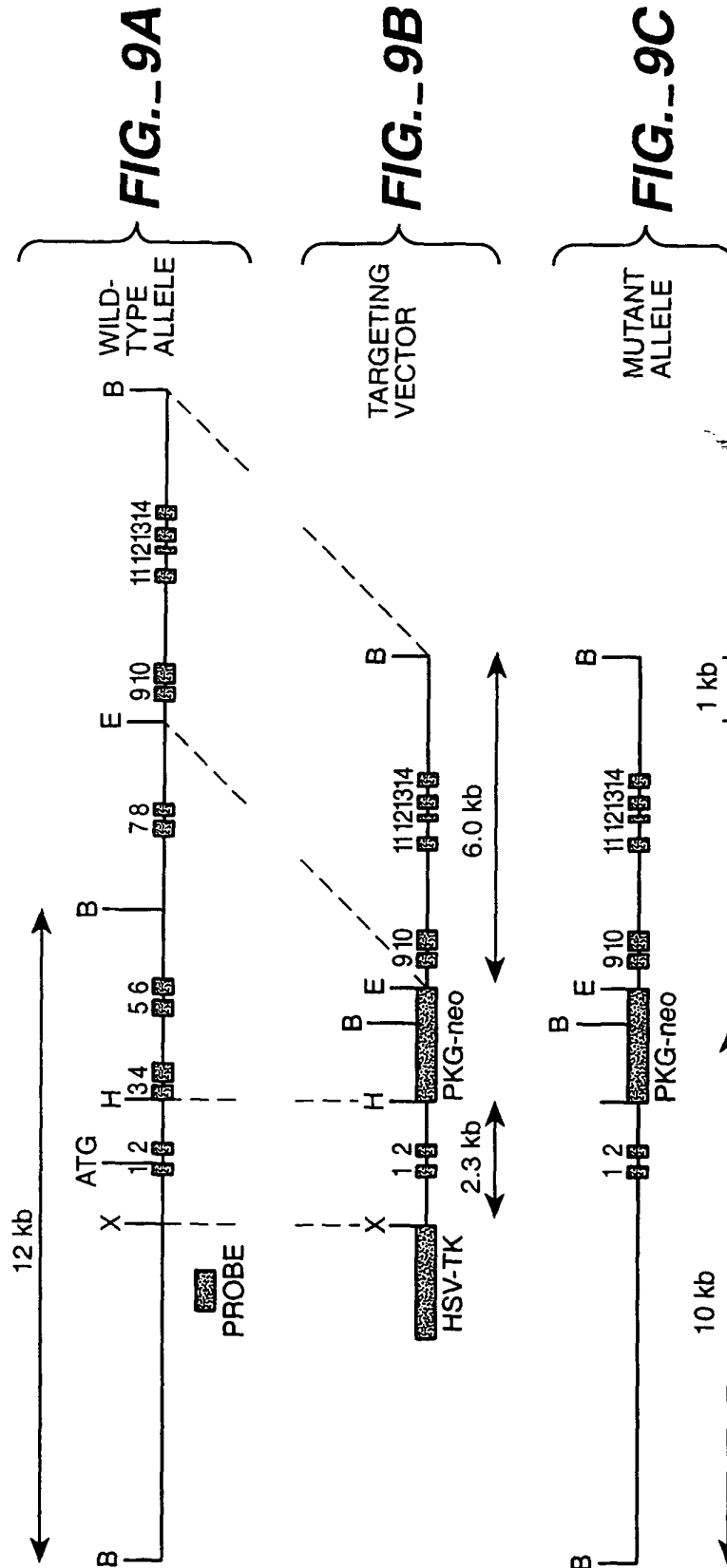
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FIG._10A

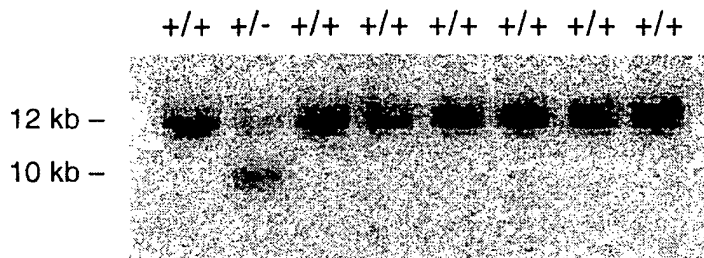


FIG._10B

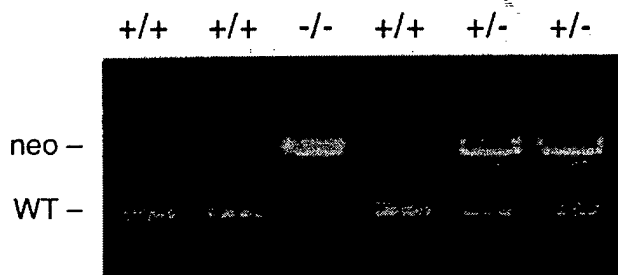
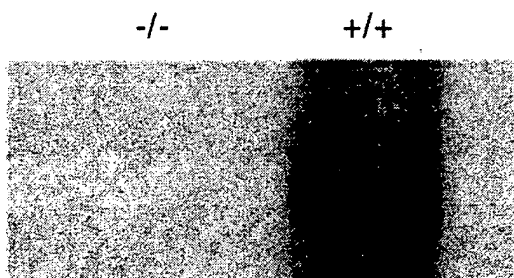
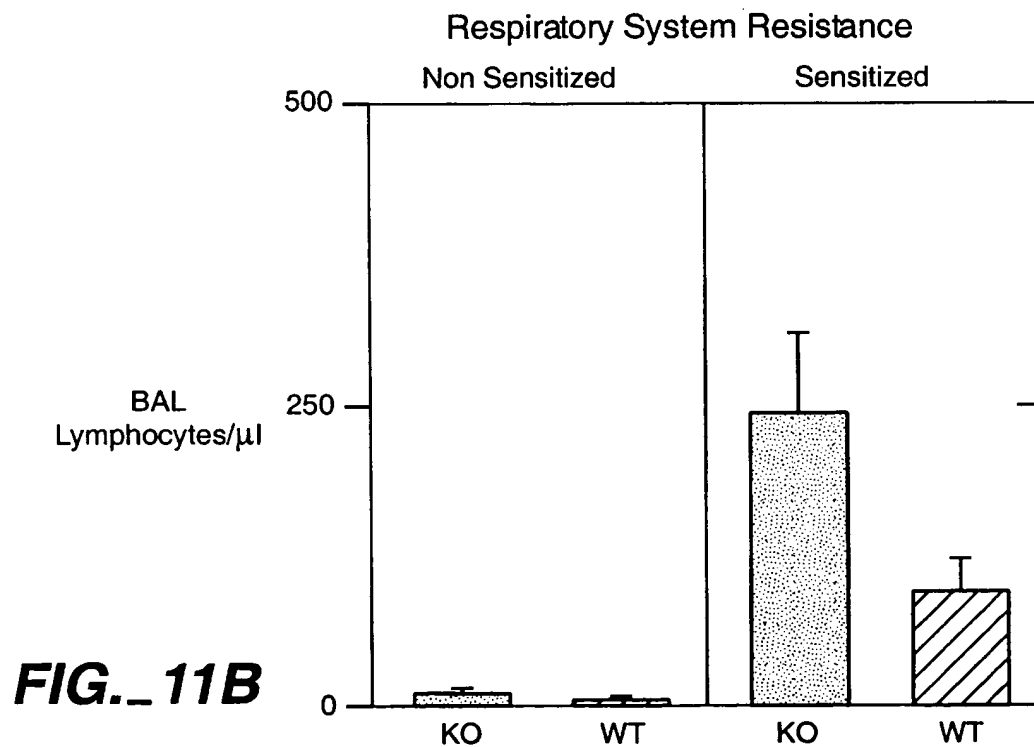
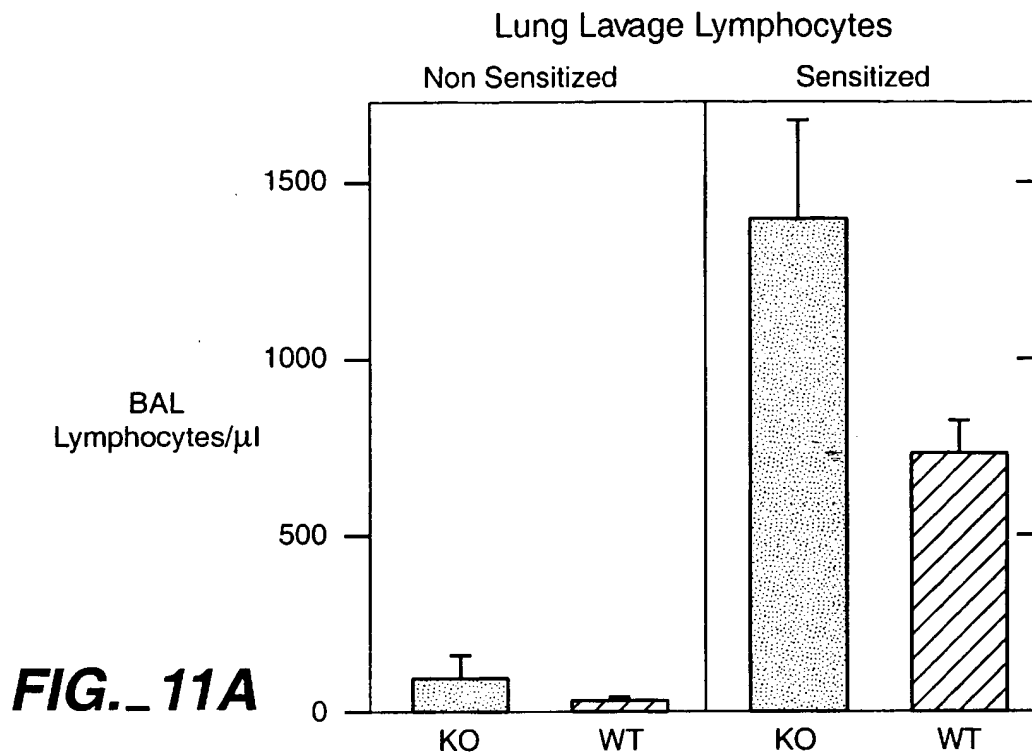


FIG._10C



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FIG. 12A

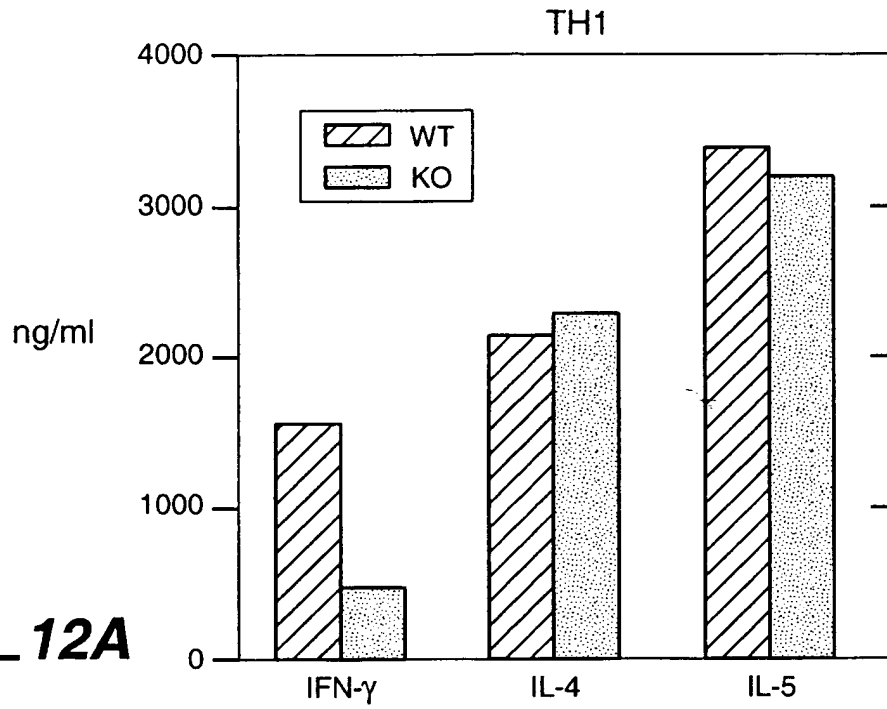
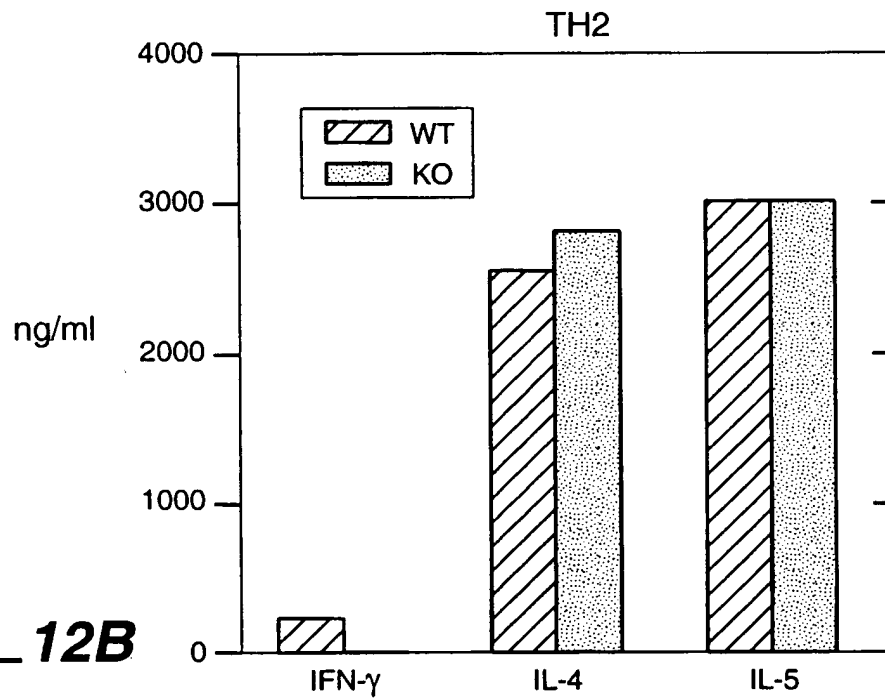


FIG. 12B



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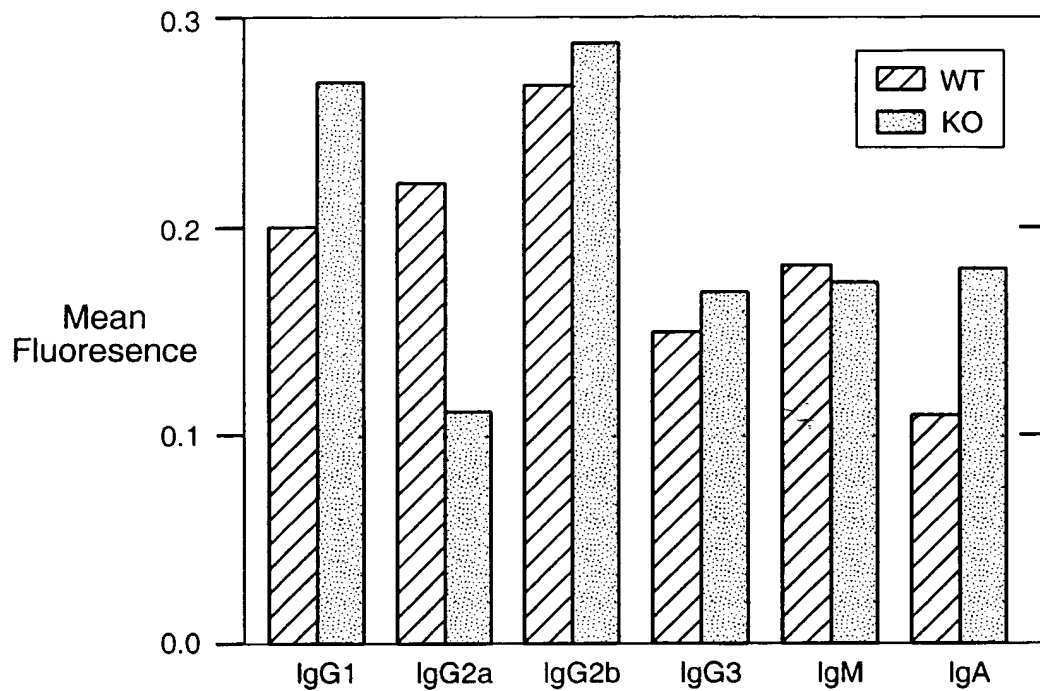


FIG._13

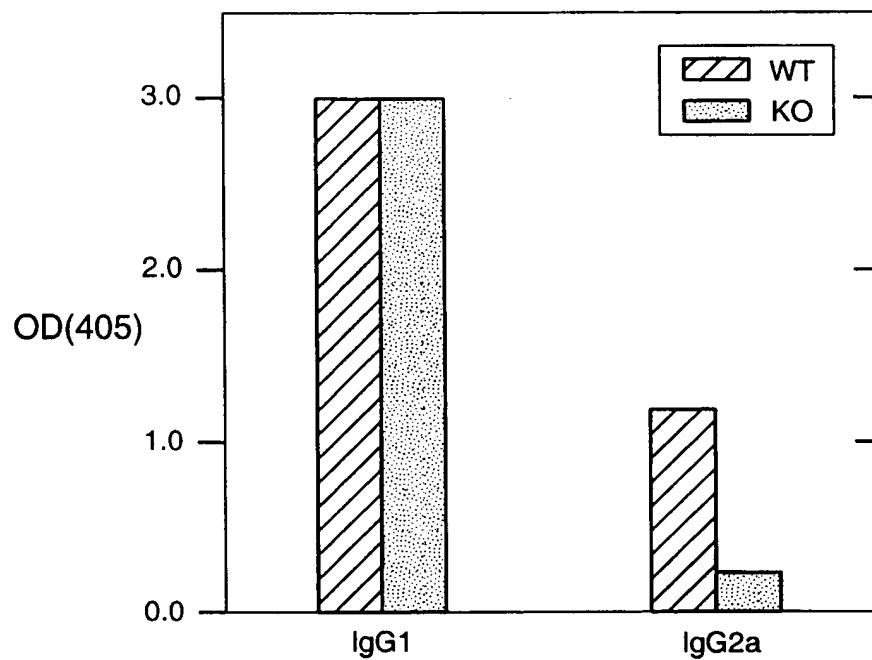


FIG._14

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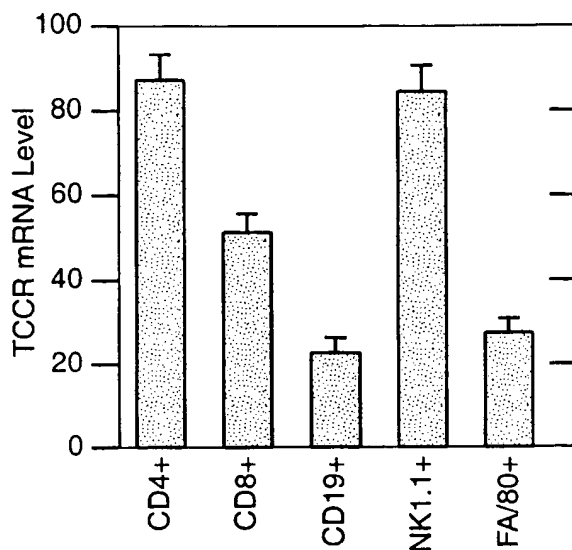


FIG._15A

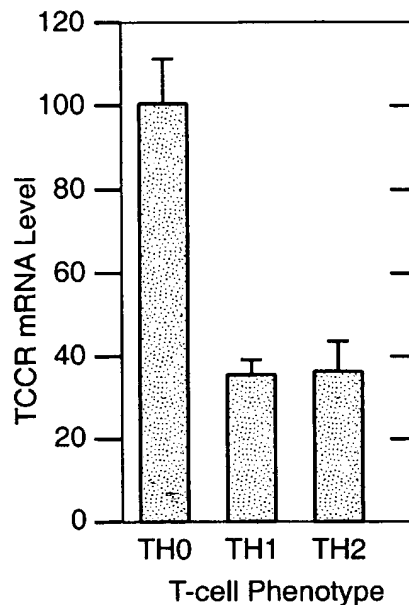


FIG._15B

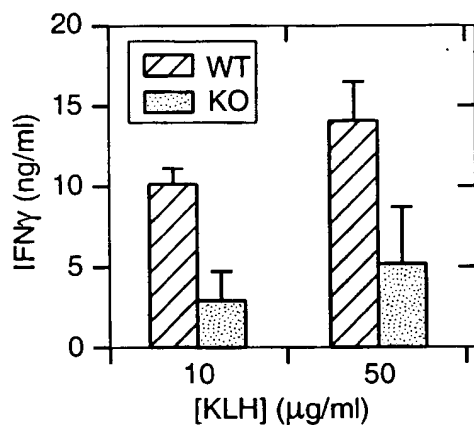


FIG._16A

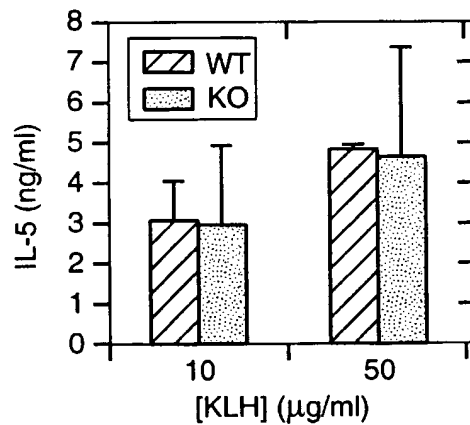


FIG._16C

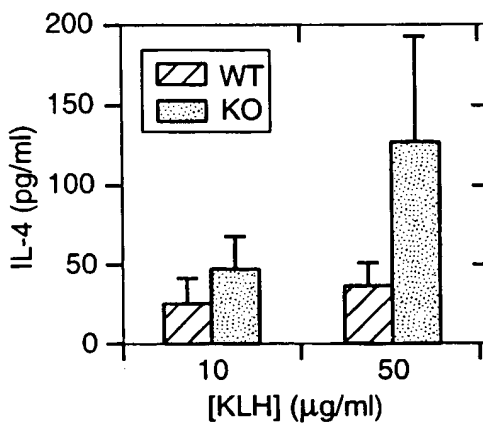


FIG._16B

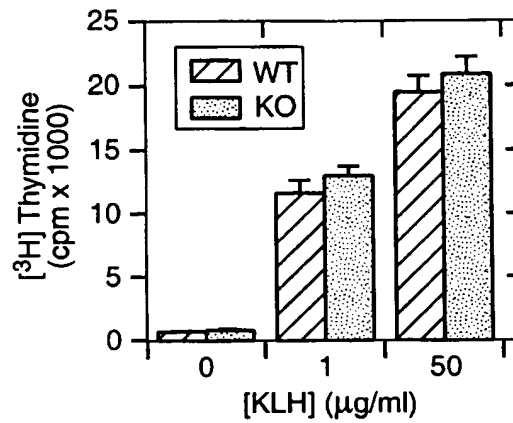
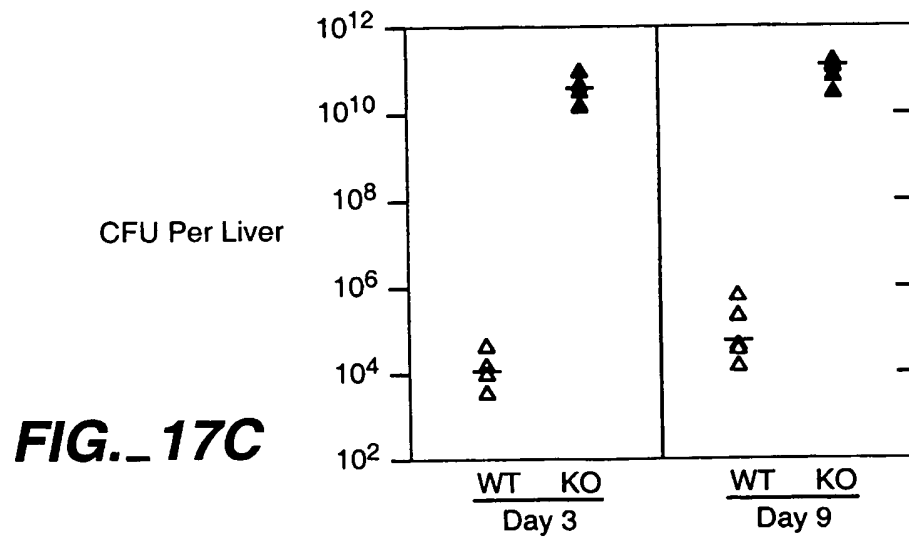
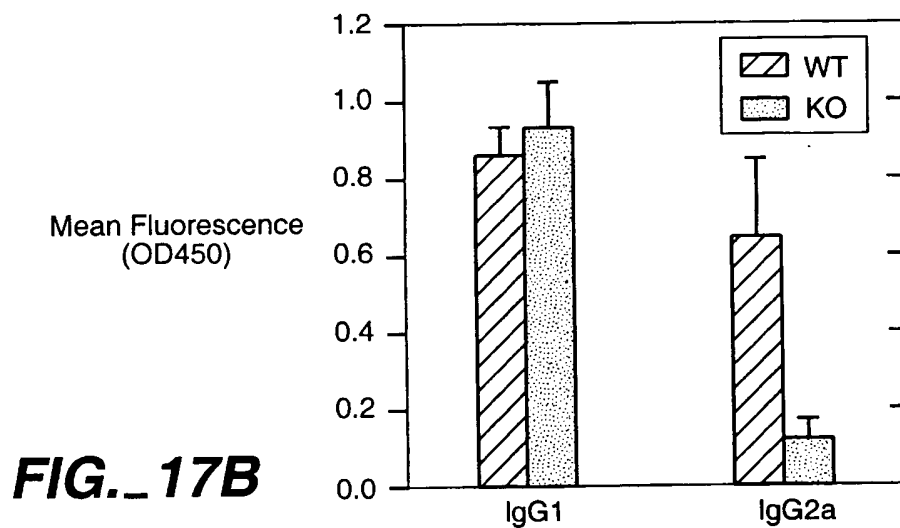
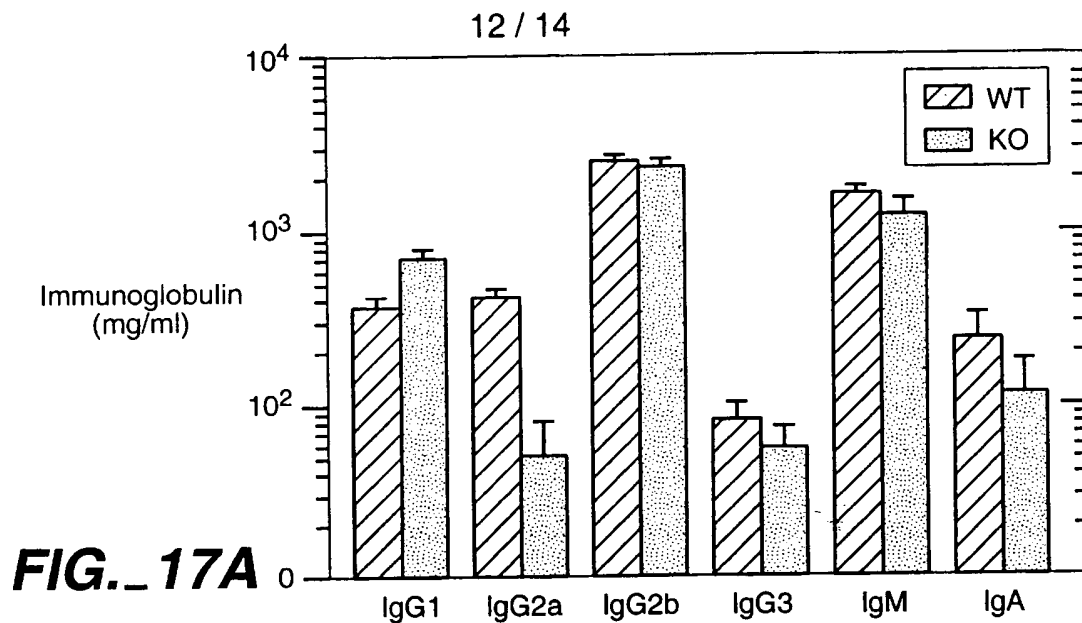


FIG._16D

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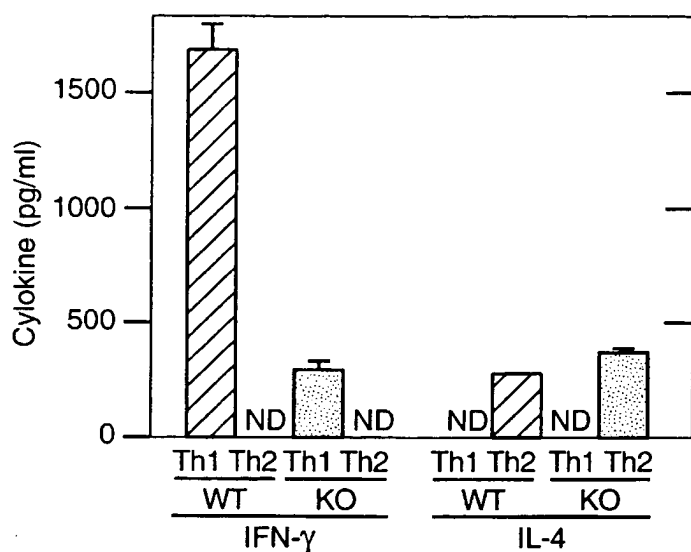


FIG. 18A

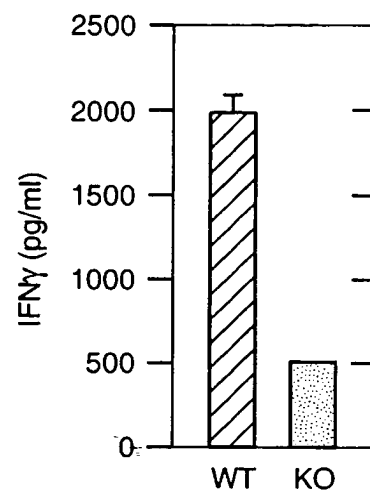


FIG. 18B

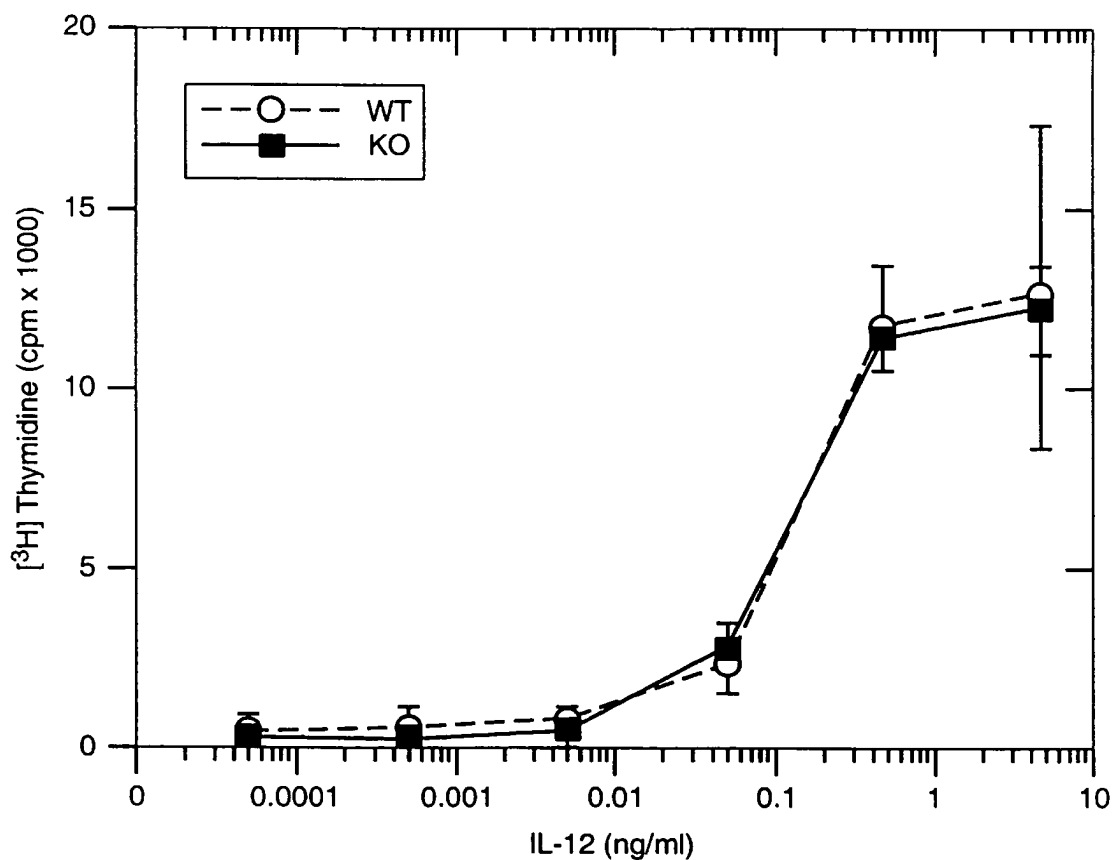
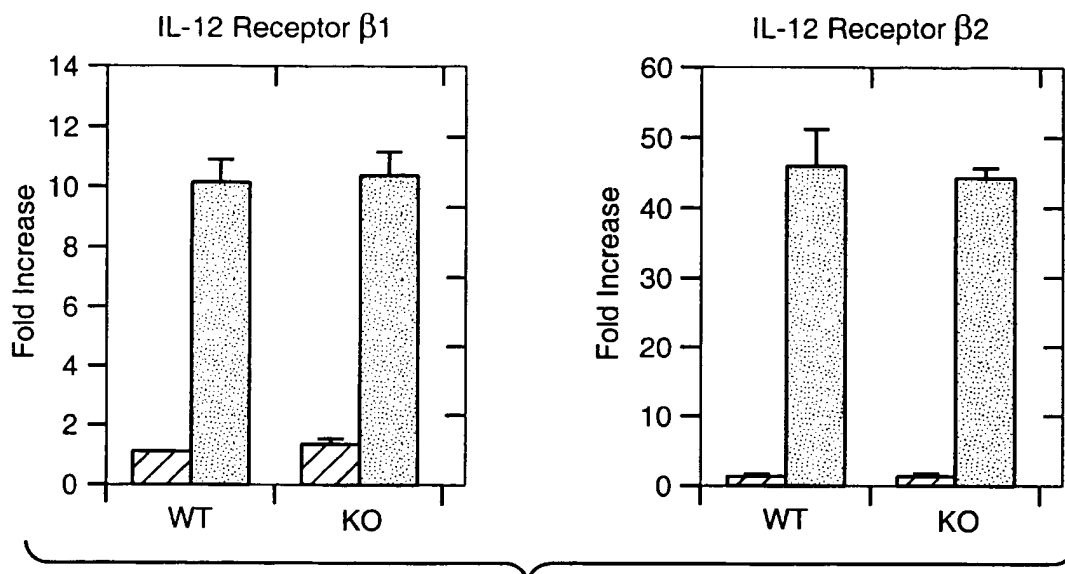


FIG. 18C

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**FIG._18D**

<u>Primer / Probe</u>	<u>Sequence</u>	<u>SEQ ID NO:</u>
mTCCR, sense, Taqman	TGGTCTCTCCTGGCAACAGC	5
mTCCR, as, Taqman	AGCCAAGCACACCAGAGACA	6
mTCCR, Taqman probe	CAGCTGGGTGCCTCCCAACAA	7
mRPL19, sense, Taqman	ATCCGCAAGCCTGTGACTGT	8
mRPL19, as, Taqman	TCGGGCCAGGGTGTTTTT	9
mRPL19, Taqman probe	TTCCCGGGCTCGTTGCCG	10
mIL12Rb1, sense, Taqman	TCGCGTCTCTGGGAAGCT	11
mIL12Rb1, as, Taqman	TTTAAGCCAATGTATCCGAGACTG	12
mIL12Rb1, Taqman probe	CGCCAGCGTCCTCCTCGTGG	13
mIL12Rb2, sense, Taqman	CAAGCATTTGCATCGCTATCA	14
mIL12Rb2, as, Taqman	AATGCCTTTTGCCGGAAGT	15
mIL12Rb2, Taqman probe	ACGAATTGAGAACGTGCCCCACCGT	16

FIG._19